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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re application of:	Group Art Unit: 1656
Xiao Bing WANG	Examiner: A. Spiegler
Serial No.: 09/618,129	
Filed: July 17, 2000	
For: **DETECTION OF SEQUENCE VARIATION OF NUCLEIC ACID BY SHIFTED TERMINATION ANALYSIS**	

AMENDMENT UNDER 37 C.F.R. §1.111

Commissioner for Patents
Washington, D.C. 20231

Sir:

In response to the Office Action dated September 20, 2001 in connection with the above-identified application, please enter and consider the following amendment and remarks:

In the Specification:

At page 9, please amend the first paragraph under "Brief Description of the Drawings":

-- FIGS. 1A-1C. A schematic drawing of a preferred embodiment of the mutation detection method of the invention is shown. "L" represents the wild-type nucleotide, which can include A, G, C, T, or U. "L*" represents an unlabeled terminator such as a dideoxynucleotide that is complementary to L. "M" represents a mutation at site L, and the mutant nucleotide can include A, G, C, T, or U. "n" represents one or multiple nucleotides or nucleotide analogues, including A, G, C, T, and U. "y" represents a nucleotide or nucleotide analogue, including A, G, C, T, or U, labeled with a detectable marker and complementary to M or n. --